

SEQUENCE LISTING

<110> Donoho, Gregory
 Scoville, John
 Turner, C. Alexander Jr.
 Friedrich, Glenn
 Zambrowicz, Brian
 Sands, Arthur T.

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Figure 1 consists of 10 subplots, labeled (a) through (j), each showing the distribution of a specific protein type across 10 fractions (A through J). The y-axis for all plots is 'Percentage of total protein' ranging from 0 to 100. The x-axis is labeled with fraction letters A through J. The legend indicates that the bars represent the percentage of total protein in each fraction for the respective protein type.

- (a) Protein A: Fractions A and B show high percentages (~80% and ~70% respectively), while others are low.
- (b) Protein B: Fractions A and B show high percentages (~80% and ~70% respectively), while others are low.
- (c) Protein C: Fractions A and B show high percentages (~80% and ~70% respectively), while others are low.
- (d) Protein D: Fractions A and B show high percentages (~80% and ~70% respectively), while others are low.
- (e) Protein E: Fractions A and B show high percentages (~80% and ~70% respectively), while others are low.
- (f) Protein F: Fractions A and B show high percentages (~80% and ~70% respectively), while others are low.
- (g) Protein G: Fractions A and B show high percentages (~80% and ~70% respectively), while others are low.
- (h) Protein H: Fractions A and B show high percentages (~80% and ~70% respectively), while others are low.
- (i) Protein I: Fractions A and B show high percentages (~80% and ~70% respectively), while others are low.
- (j) Protein J: Fractions A and B show high percentages (~80% and ~70% respectively), while others are low.

[illegible][illegible][illegible][illegible]

Figure 1 consists of 11 line graphs, labeled (a) through (k), arranged vertically. Each graph plots 'Number of children' on the y-axis against 'Number of children' on the x-axis. The data points are connected by lines, showing a general upward trend across all variables. The variables represented are: (a) Number of children, (b) Number of children, (c) Number of children, (d) Number of children, (e) Number of children, (f) Number of children, (g) Number of children, (h) Number of children, (i) Number of children, (j) Number of children, and (k) Number of children. The graphs show that as the number of children increases, the values for these variables also tend to increase, with some variables showing a more significant increase than others.

Figure 1 consists of 12 line graphs arranged in a 3x4 grid. Each graph plots the 'Rate of change of number of cells in mitotic pool' on the y-axis against a specific factor on the x-axis. The factors are: (a) Time, (b) Cell cycle time, (c) Cell cycle length, (d) Cell cycle duration, (e) Cell cycle frequency, (f) Cell cycle rate, (g) Cell cycle time, (h) Cell cycle length, (i) Cell cycle duration, (j) Cell cycle frequency, (k) Cell cycle rate, and (l) Cell cycle time. The graphs show that the rate of change is highest when the cell cycle time is short and decreases as the cell cycle time increases.

Figure 1 consists of 11 line graphs, labeled (a) through (k), arranged vertically. Each graph plots 'Number of children' on the y-axis against 'Number of children' on the x-axis. The data points are connected by lines, showing a general upward trend across all variables. The variables represented are: (a) Number of children, (b) Number of children, (c) Number of children, (d) Number of children, (e) Number of children, (f) Number of children, (g) Number of children, (h) Number of children, (i) Number of children, (j) Number of children, and (k) Number of children. The graphs show that as the number of children increases, the values for these variables also tend to increase, with some variables showing a more significant increase than others.

Figure 1 consists of 11 sub-graphs, labeled (a) through (k), each showing the percentage of total catch for a specific fish species over time from 1970 to 1990. The y-axis for all graphs is 'Percentage of total catch' ranging from 0 to 100. The x-axis is 'Year' from 1970 to 1990. Three scenarios are compared: (a) no trawling (solid line), (b) trawling in the 1970s (dashed line), and (c) trawling in the 1980s (dotted line). The species are: (a) Atlantic cod, (b) Atlantic herring, (c) Atlantic mackerel, (d) Atlantic plaice, (e) Atlantic sandeel, (f) Atlantic whiting, (g) European eel, (h) European hake, (i) European plaice, (j) European sandeel, and (k) European whiting. In general, the 'no trawling' scenario maintains a high percentage of catch (near 100%) for most species. The 'trawling in the 1970s' scenario shows a significant decline in catch percentage for several species, particularly in the 1980s. The 'trawling in the 1980s' scenario shows the most severe declines, with many species dropping to near zero by the end of the period.

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 <212> PRT
 <213> Homo sapiens

<400> 22
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 His Gln Thr Gly Ala Arg Trp Arg Pro Leu Pro Gln Arg Glu Ser Gln
 20 25 30
 Gly Leu Met Gly Gly Asn Gly Arg Gly Thr Phe Thr Asp Arg Lys Ala
 35 40 45
 Gln Pro Gly Asp Phe Leu Gly Leu Leu Ala Arg Gly Thr Thr Pro Ser
 50 55 60
 Pro Thr Thr Ala Ala Pro Ser Ser Arg Cys Trp Phe Arg Glu Gly Thr
 65 70 75 80
 Thr Met Tyr Ala Leu Tyr Ile Thr Val His Gly Tyr Phe Leu Ile Thr
 85 90 95
 Phe Leu Phe Gly Met Val Val Leu Ala Leu Val Val Trp Lys Ile Phe
 100 105 110
 Thr Leu Ser Arg Ala Thr Ala Val Lys Glu Arg Gly Lys Asn Arg Cys
 115 120 125
 Ser Pro Cys Trp Ala Ser Arg Ala Leu Gln Val Gly Cys Pro Ser Ser
 130 135 140
 Ile Ser Gly Pro Ile Ser Cys Asp Gln Lys Gly Arg Ile Met
 145 150 155

<210> 23
 <211> 1566
 <212> DNA
 <213> Homo sapiens

<400> 23
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 caggaaaagc ccaccgaagg gccaaagaac acctgcctgg ggagcaaca catgtacgac 120

Lys Leu Ala Glu Pro Leu Glu Ile Val Phe Ser His Gln Arg Pro Pro
 195 200 205
 Pro Asn Met Thr Leu Thr Cys Val Phe Trp Asp Val Thr Lys Gly Thr
 210 215 220
 Thr Gly Asp Trp Ser Ser Glu Gly Cys Ser Thr Glu Val Arg Pro Glu
 225 230 235 240
 Gly Thr Val Cys Cys Cys Asp His Leu Thr Phe Phe Ala Leu Leu Leu
 245 250 255
 Arg Pro Thr Leu Asp Gln Ser Thr Val His Ile Leu Thr Arg Ile Ser
 260 265 270
 Gln Ala Gly Cys Gly Val Ser Met Ile Phe Leu Ala Phe Thr Ile Ile
 275 280 285
 Leu Tyr Ala Phe Leu Arg Leu Ser Arg Glu Arg Phe Lys Ser Glu Asp
 290 295 300
 Ala Pro Lys Ile His Val Ala Leu Gly Gly Ser Leu Phe Leu Leu Asn
 305 310 315 320
 Leu Ala Phe Leu Val Asn Val Gly Ser Gly Ser Lys Gly Ser Asp Ala
 325 330 335
 Ala Cys Trp Ala Arg Gly Ala Val Phe His Tyr Phe Leu Leu Cys Ala
 340 345 350
 Phe Thr Trp Met Gly Leu Glu Ala Phe His Leu Tyr Leu Leu Ala Val
 355 360 365
 Arg Val Phe Asn Thr Tyr Phe Gly His Tyr Phe Leu Lys Leu Ser Leu
 370 375 380
 Val Gly Trp Gly Leu Pro Ala Leu Met Val Ile Gly Thr Gly Ser Ala
 385 390 395 400
 Asn Ser Tyr Gly Leu Tyr Thr Ile Arg Asp Arg Glu Asn Arg Thr Ser
 405 410 415
 Leu Glu Leu Cys Trp Phe Arg Glu Gly Thr Thr Met Tyr Ala Leu Tyr
 420 425 430
 Ile Thr Val His Gly Tyr Phe Leu Ile Thr Phe Leu Phe Gly Met Val
 435 440 445
 Val Leu Ala Leu Val Val Trp Lys Ile Phe Thr Leu Ser Arg Ala Thr
 450 455 460
 Ala Val Lys Glu Arg Gly Lys Asn Arg Lys Lys Val Leu Thr Leu Leu
 465 470 475 480
 Gly Leu Ser Ser Leu Val Gly Val Thr Trp Gly Leu Ala Ile Phe Thr
 485 490 495
 Pro Leu Gly Leu Ser Thr Val Tyr Ile Phe Ala Leu Phe Asn Ser Leu
 500 505 510
 Gln Gly Glu Ala Pro Ala Pro Gly Arg
 515 520

<210> 25
 <211> 936
 <212> DNA
 <213> Homo sapiens

<400> 25
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 gagggctgct ccacggaggt cagacctgag gggaccgtgt gctgctgtga ccacctgacc 120
 tttttcgccc tgctcctgag acccaccttg gaccagtcca cggtgcatat cctcacacgc 180
 atctcccagg cgggctgtgg ggtctccatg atcttctctg ccttcaccat tattctttat 240
 gcctttctga ggctttccc ggagagggtc aagtcagaag atgccccaaa gatccacgtg 300
 gccctgggtg gcagcctggt cctcctgaat ctggccttct tggccaatgt ggggagtggc 360
 tcaaaggggt ctgatgctgc ctgctgggccc cggggggctg tcttcacta cttcctgctc 420

tgtgccttca	cctggatggg	ccttgaagcc	ttccacctct	acctgctcgc	tgctcagggtc	480
ttcaacacct	acttcgggca	ctacttctctg	aagctgagcc	tggtgggctg	gggcctgccc	540
gccctgatgg	tcacgcggcac	tgggagtgcc	aacagctacg	gcctctacac	catccgtgat	600
agggagaacc	gcacctctct	ggagctatgc	tggttccgtg	aagggaacaac	catgtacgcc	660
ctctatatca	cgcgccacgg	ctacttcctc	atcaccttcc	tctttggcat	ggtggctcctg	720
gccctgggtg	tctggaagat	cttcaccctg	tcccgtgcta	cagcgggtcaa	ggagcggggg	780
aagaaccgga	agaagggtgct	caccctgctg	ggcctctcga	gcctgggtggg	tgtagacatgg	840
gggttggcca	tcttcacccc	gttgggcctc	tccaccgtct	acatctttgc	acttttcaac	900
tccttgcaag	gtgaggcccc	tgcaccaggg	aggtga			936

<210> 26
 <211> 311
 <212> PRT
 <213> Homo sapiens

<400> 26

Met	Thr	Leu	Thr	Cys	Val	Phe	Trp	Asp	Val	Thr	Lys	Gly	Thr	Thr	Gly	
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Asp	Trp	Ser	Ser	Glu	Gly	Cys	Ser	Thr	Glu	Val	Arg	Pro	Glu	Gly	Thr	
			20					25					30			
Val	Cys	Cys	Cys	Asp	His	Leu	Thr	Phe	Phe	Ala	Leu	Leu	Leu	Arg	Pro	
			35				40					45				
Thr	Leu	Asp	Gln	Ser	Thr	Val	His	Ile	Leu	Thr	Arg	Ile	Ser	Gln	Ala	
	50					55					60					
Gly	Cys	Gly	Val	Ser	Met	Ile	Phe	Leu	Ala	Phe	Thr	Ile	Ile	Leu	Tyr	
65				70					75					80		
Ala	Phe	Leu	Arg	Leu	Ser	Arg	Glu	Arg	Phe	Lys	Ser	Glu	Asp	Ala	Pro	
			85					90					95			
Lys	Ile	His	Val	Ala	Leu	Gly	Gly	Ser	Leu	Phe	Leu	Leu	Asn	Leu	Ala	
			100				105						110			
Phe	Leu	Val	Asn	Val	Gly	Ser	Gly	Ser	Lys	Gly	Ser	Asp	Ala	Ala	Cys	
		115				120						125				
Trp	Ala	Arg	Gly	Ala	Val	Phe	His	Tyr	Phe	Leu	Leu	Cys	Ala	Phe	Thr	
	130				135						140					
Trp	Met	Gly	Leu	Glu	Ala	Phe	His	Leu	Tyr	Leu	Leu	Ala	Val	Arg	Val	
145				150					155					160		
Phe	Asn	Thr	Tyr	Phe	Gly	His	Tyr	Phe	Leu	Lys	Leu	Ser	Leu	Val	Gly	
			165					170					175			
Trp	Gly	Leu	Pro	Ala	Leu	Met	Val	Ile	Gly	Thr	Gly	Ser	Ala	Asn	Ser	
		180					185						190			
Tyr	Gly	Leu	Tyr	Thr	Ile	Arg	Asp	Arg	Glu	Asn	Arg	Thr	Ser	Leu	Glu	
	195					200						205				
Leu	Cys	Trp	Phe	Arg	Glu	Gly	Thr	Thr	Met	Tyr	Ala	Leu	Tyr	Ile	Thr	
	210					215					220					
Val	His	Gly	Tyr	Phe	Leu	Ile	Thr	Phe	Leu	Phe	Gly	Met	Val	Val	Leu	
225					230				235					240		
Ala	Leu	Val	Val	Trp	Lys	Ile	Phe	Thr	Leu	Ser	Arg	Ala	Thr	Ala	Val	
			245					250					255			
Lys	Glu	Arg	Gly	Lys	Asn	Arg	Lys	Lys	Val	Leu	Thr	Leu	Leu	Gly	Leu	
			260				265						270			
Ser	Ser	Leu	Val	Gly	Val	Thr	Trp	Gly	Leu	Ala	Ile	Phe	Thr	Pro	Leu	
	275					280					285					
Gly	Leu	Ser	Thr	Val	Tyr	Ile	Phe	Ala	Leu	Phe	Asn	Ser	Leu	Gln	Gly	
	290				295						300					
Glu	Ala	Pro	Ala	Pro	Gly	Arg										
305					310											

[illegible]

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<210> 28
<211> 372
<212> PRT
<213> Homo sapiens
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18

[illegible]

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<210> 31
<211> 528
<212> DNA
<213> Homo sapiens
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<400> 31						
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gccaggtggc	ggccccctcc	acagcgggag	agccagggat	tgatgggttg	aaatgggaga	120
ggcaccttca	cagacagaaa	agctcagcca	ggggacttcc	tgggtttgct	ggccagaggt	180
accactccca	gtcccaccac	agctgcccc	tcctccagat	gctggttccg	tgaagggaca	240
accatgtacg	ccctctatat	caccgtccac	ggctacttcc	tcatcacctt	cctctttggc	300
atggtgggtc	tggccctggt	ggtctggaag	atcttcaccc	tgccccgtgc	tacagcggtc	360
aaggagcggg	ggaagaaccg	gaagaagggt	ctcacccctg	tgggcctctc	gagcctgggt	420
ggtgtgacat	gggggttggc	catcttcacc	cgttgggcc	tctccaccgt	ctacatcttt	480
gcacttttca	actccttgca	aggtgaggcc	cctgcaccag	ggaggtga		528

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<210> 32
<211> 175
<212> PRT
<213> Homo sapiens
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<400> 32															
Met	Gly	Gln	Met	Lys	His	Val	Phe	Glu	Val	Thr	Leu	Ala	Leu	Lys	Arg
1				5					10					15	
His	Gln	Thr	Gly	Ala	Arg	Trp	Arg	Pro	Leu	Pro	Gln	Arg	Glu	Ser	Gln
			20					25					30		
Gly	Leu	Met	Gly	Gly	Asn	Gly	Arg	Gly	Thr	Phe	Thr	Asp	Arg	Lys	Ala
		35					40					45			

Gln	Pro	Gly	Asp	Phe	Leu	Gly	Leu	Leu	Ala	Arg	Gly	Thr	Thr	Pro	Ser
50						55					60				
Pro	Thr	Thr	Ala	Ala	Pro	Ser	Ser	Arg	Cys	Trp	Phe	Arg	Glu	Gly	Thr
65					70				75					80	
Thr	Met	Tyr	Ala	Leu	Tyr	Ile	Thr	Val	His	Gly	Tyr	Phe	Leu	Ile	Thr
			85					90						95	
Phe	Leu	Phe	Gly	Met	Val	Val	Leu	Ala	Leu	Val	Val	Trp	Lys	Ile	Phe
			100					105					110		
Thr	Leu	Ser	Arg	Ala	Thr	Ala	Val	Lys	Glu	Arg	Gly	Lys	Asn	Arg	Lys
		115				120					125				
Lys	Val	Leu	Thr	Leu	Leu	Gly	Leu	Ser	Ser	Leu	Val	Gly	Val	Thr	Trp
	130					135					140				
Gly	Leu	Ala	Ile	Phe	Thr	Pro	Leu	Gly	Leu	Ser	Thr	Val	Tyr	Ile	Phe
145					150				155					160	
Ala	Leu	Phe	Asn	Ser	Leu	Gln	Gly	Glu	Ala	Pro	Ala	Pro	Gly	Arg	
			165					170						175	

<210> 33
 <211> 1458
 <212> DNA
 <213> Homo sapiens

<400> 33
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 atcttcaact tgaatgacaa ggctttgtgc ttcaccaagt gcaggcagtc gggcagcgac 180
 tctgcaatg tggaaaactt gcagagatac tggctaaact acgaggccca tctgatgaag 240
 gaaggtttga cgcagaaggt gaacacgcct ttctgaagg ctttggtcca gaacctcagc 300
 accaactctg cagaagactt ctatttctct ctggagccct ctcaggttcc gaggcaggtg 360
 atgaaggacg aggacaagcc ccctgacaga gtgcgacttc ccaagagcct ttttcgatcc 420
 ctgccaggca acaggtctgt ggtccgcttg gccgtcacca ttctggacat tgggtccaggg 480
 actctcttca agggcccccg gctcggcctg ggagatggca gggcgctgtt gaacaatcgc 540
 ctggtgggtt tgagtgtggg acaaatgcat gtcaccaagc tggctgagcc tctggagatc 600
 gtcttctctc accagcgacc gcccctaacc atgacctca cctgtgtatt ctgggatgtg 660
 actaaaggga cactggaga ctggtcttct gagggctgct ccacggaggt cagacctgag 720
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 gaccagtcca cgggtcatat cctcacacgc atctcccagg cgggctgtgg ggtctccatg 840
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 aagtcagaag atgccccaaa gatccacgtg gccctgggtg gcagcctgtt cctcctgaat 960
 ctggccttct tgggtcaatgt ggggagtggc tcaaaggggt ctgatgctgc ctgctgggcc 1020
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 ttccacctct acctgctcgc tgtcagggtc ttcaaacact acttcgggca ctacttcctg 1140
 aagctgagcc tgggtgggctg gggcctgccc gccctgatgg tcatcggcac tgggagtgcc 1200
 aacagctacg gcctctacac catccgtgat agggagaacc gcacctctct ggagctatgc 1260
 tggttccgtg aagggacaac catgtacgcc ctctatatca ccgtccacgg ctacttcctc 1320
 atcaccttcc tctttggcat ggtggtcctg gccctgggtg tctggaagat cttcacctctg 1380
 tcccgtgcta cagcgggtcaa ggagcggggg aagaaccggt gtcaccctg ctgggcctct 1440
 cgagcctggt ggggtgtga 1458

<210> 34
 <211> 485
 <212> PRT
 <213> Homo sapiens

<400> 34
 Met Ala Thr Pro Arg Gly Leu Gly Ala Leu Leu Leu Leu Leu Leu

1	5						10						15			
Pro	Thr	Ser	Gly	Gln	Glu	Lys	Pro	Thr	Glu	Gly	Pro	Arg	Asn	Thr	Cys	
			20					25					30			
Leu	Gly	Ser	Asn	Asn	Met	Tyr	Asp	Ile	Phe	Asn	Leu	Asn	Asp	Lys	Ala	
		35					40					45				
Leu	Cys	Phe	Thr	Lys	Cys	Arg	Gln	Ser	Gly	Ser	Asp	Ser	Cys	Asn	Val	
	50					55					60					
Glu	Asn	Leu	Gln	Arg	Tyr	Trp	Leu	Asn	Tyr	Glu	Ala	His	Leu	Met	Lys	
65					70					75					80	
Glu	Gly	Leu	Thr	Gln	Lys	Val	Asn	Thr	Pro	Phe	Leu	Lys	Ala	Leu	Val	
				85					90					95		
Gln	Asn	Leu	Ser	Thr	Asn	Thr	Ala	Glu	Asp	Phe	Tyr	Phe	Ser	Leu	Glu	
			100					105					110			
Pro	Ser	Gln	Val	Pro	Arg	Gln	Val	Met	Lys	Asp	Glu	Asp	Lys	Pro	Pro	
		115					120					125				
Asp	Arg	Val	Arg	Leu	Pro	Lys	Ser	Leu	Phe	Arg	Ser	Leu	Pro	Gly	Asn	
	130					135					140					
Arg	Ser	Val	Val	Arg	Leu	Ala	Val	Thr	Ile	Leu	Asp	Ile	Gly	Pro	Gly	
145					150					155					160	
Thr	Leu	Phe	Lys	Gly	Pro	Arg	Leu	Gly	Leu	Gly	Asp	Gly	Ser	Gly	Val	
				165					170					175		
Leu	Asn	Asn	Arg	Leu	Val	Gly	Leu	Ser	Val	Gly	Gln	Met	His	Val	Thr	
			180					185					190			
Lys	Leu	Ala	Glu	Pro	Leu	Glu	Ile	Val	Phe	Ser	His	Gln	Arg	Pro	Pro	
		195					200					205				
Pro	Asn	Met	Thr	Leu	Thr	Cys	Val	Phe	Trp	Asp	Val	Thr	Lys	Gly	Thr	
	210					215					220					
Thr	Gly	Asp	Trp	Ser	Ser	Glu	Gly	Cys	Ser	Thr	Glu	Val	Arg	Pro	Glu	
225					230					235					240	
Gly	Thr	Val	Cys	Cys	Cys	Asp	His	Leu	Thr	Phe	Phe	Ala	Leu	Leu	Leu	
				245					250					255		
Arg	Pro	Thr	Leu	Asp	Gln	Ser	Thr	Val	His	Ile	Leu	Thr	Arg	Ile	Ser	
			260					265					270			
Gln	Ala	Gly	Cys	Gly	Val	Ser	Met	Ile	Phe	Leu	Ala	Phe	Thr	Ile	Ile	
		275					280					285				
Leu	Tyr	Ala	Phe	Leu	Arg	Leu	Ser	Arg	Glu	Arg	Phe	Lys	Ser	Glu	Asp	
	290					295					300					
Ala	Pro	Lys	Ile	His	Val	Ala	Leu	Gly	Gly	Ser	Leu	Phe	Leu	Leu	Asn	
305					310					315					320	
Leu	Ala	Phe	Leu	Val	Asn	Val	Gly	Ser	Gly	Ser	Lys	Gly	Ser	Asp	Ala	
				325					330					335		
Ala	Cys	Trp	Ala	Arg	Gly	Ala	Val	Phe	His	Tyr	Phe	Leu	Leu	Cys	Ala	
			340					345					350			
Phe	Thr	Trp	Met	Gly	Leu	Glu	Ala	Phe	His	Leu	Tyr	Leu	Leu	Ala	Val	
		355					360					365				

450 455 460
 Ala Val Lys Glu Arg Gly Lys Asn Arg Cys Ser Pro Cys Trp Ala Ser
 465 470 475 480
 Arg Ala Trp Trp Val
 485

<210> 35
 <211> 828
 <212> DNA
 <213> Homo sapiens

<400> 35
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 gagggctgct ccacggaggt cagacctgag gggacctgtg gctgctgtga ccacctgacc 120
 tttttcgccc tgctcctgag acccaccttg gaccagtcca cgggtgcata cctcacacgc 180
 atctcccagg cgggctgtgg ggtctccatg atcttcctgg ccttcaccat tattctttat 240
 gcctttctga ggctttcccg ggagaggttc aagtcagaag atgccccaaa gatccacgtg 300
 gccctgggtg gcagcctggt cctcctgaat ctggccttct tgggtcaatgt ggggagtggc 360
 tcaaaggggt ctgatgctgc ctgctgggcc cggggggctg tcttcacta cttcctgctc 420
 tgtgccttca cctggatggg ccttgaagcc ttccacctct acctgctcgc tgtcagggtc 480
 ttcaacacct acttcgggca ctacttctg aagctgagcc tgggtgggctg gggcctgccc 540
 gccctgatgg tcatcggcac tgggagtgcc aacagctacg gcctctacac catccgtgat 600
 agggagaacc gcacctctct ggagctatgc tgggtccgtg aagggaacaac catgtacgcc 660
 ctctatatca ccgtccacgg ctacttcctc atcaccttcc tctttggcat ggtggctcgt 720
 gccctgggtg tctggaagat cttcacctct tcccgtgcta cagcggtcaa ggagcggggg 780
 aagaaccggt gtcaccctg ctgggcctct cgagcctggt ggggtgtga 828

<210> 36
 <211> 275
 <212> PRT
 <213> Homo sapiens

<400> 36
 Met Thr Leu Thr Cys Val Phe Trp Asp Val Thr Lys Gly Thr Thr Gly
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 Asp Trp Ser Ser Glu Gly Cys Ser Thr Glu Val Arg Pro Glu Gly Thr
 20 25 30
 Val Cys Cys Cys Asp His Leu Thr Phe Phe Ala Leu Leu Leu Arg Pro
 35 40 45
 Thr Leu Asp Gln Ser Thr Val His Ile Leu Thr Arg Ile Ser Gln Ala
 50 55 60
 Gly Cys Gly Val Ser Met Ile Phe Leu Ala Phe Thr Ile Ile Leu Tyr
 65 70 75 80
 Ala Phe Leu Arg Leu Ser Arg Glu Arg Phe Lys Ser Glu Asp Ala Pro
 85 90 95
 Lys Ile His Val Ala Leu Gly Gly Ser Leu Phe Leu Leu Asn Leu Ala
 100 105 110
 Phe Leu Val Asn Val Gly Ser Gly Ser Lys Gly Ser Asp Ala Ala Cys
 115 120 125
 Trp Ala Arg Gly Ala Val Phe His Tyr Phe Leu Leu Cys Ala Phe Thr
 130 135 140
 Trp Met Gly Leu Glu Ala Phe His Leu Tyr Leu Leu Ala Val Arg Val
 145 150 155 160
 Phe Asn Thr Tyr Phe Gly His Tyr Phe Leu Lys Leu Ser Leu Val Gly
 165 170 175
 Trp Gly Leu Pro Ala Leu Met Val Ile Gly Thr Gly Ser Ala Asn Ser

1	5	10	15
His Pro Arg Leu Ser Arg Glu Arg Phe Lys Ser Glu Asp Ala Pro Lys			
20	25	30	
Ile His Val Ala Leu Gly Gly Ser Leu Phe Leu Leu Asn Leu Ala Phe			
35	40	45	
Leu Val Asn Val Gly Ser Gly Ser Lys Gly Ser Asp Ala Ala Cys Trp			
50	55	60	
Ala Arg Gly Ala Val Phe His Tyr Phe Leu Leu Cys Ala Phe Thr Trp			
65	70	75	80
Met Gly Leu Glu Ala Phe His Leu Tyr Leu Leu Ala Val Arg Val Phe			
85	90	95	
Asn Thr Tyr Phe Gly His Tyr Phe Leu Lys Leu Ser Leu Val Gly Trp			
100	105	110	
Gly Leu Pro Ala Leu Met Val Ile Gly Thr Gly Ser Ala Asn Ser Tyr			
115	120	125	
Gly Leu Tyr Thr Ile Arg Asp Arg Glu Asn Arg Thr Ser Leu Glu Leu			
130	135	140	
Cys Trp Phe Arg Glu Gly Thr Thr Met Tyr Ala Leu Tyr Ile Thr Val			
145	150	155	160
His Gly Tyr Phe Leu Ile Thr Phe Leu Phe Gly Met Val Val Leu Ala			
165	170	175	
Leu Val Val Trp Lys Ile Phe Thr Leu Ser Arg Ala Thr Ala Val Lys			
180	185	190	
Glu Arg Gly Lys Asn Arg Cys Ser Pro Cys Trp Ala Ser Arg Ala Trp			
195	200	205	
Trp Val			
210			

<210> 41
 <211> 420
 <212> DNA
 <213> Homo sapiens

<400> 41
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 gccagggtggc ggcccctccc acagcgggag agccagggat tgatgggtgg aaatgggaga 120
 ggcaccttca cagacagaaa agctcagcca ggggacttcc tgggtttgct ggccagaggt 180
 accactccca gtcccaccac agctgcccc tctccagat gctggttccg tgaagggaca 240
 accatgtacg ccctctatat caccgtccac ggctacttcc tcatcacctt cctctttggc 300
 atggtgggtcc tggccctggt ggtctggaag atcttcaccc tgtcccgtgc tacagcggtc 360
 aaggagcggg ggaagaaccg gtgctcaccc tgctgggcct ctcgagcctg gtgggtgtga 420

<210> 42
 <211> 139
 <212> PRT
 <213> Homo sapiens

<400> 42
 Met Gly Gln Met Lys His Val Phe Glu Val Thr Leu Ala Leu Lys Arg
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 His Gln Thr Gly Ala Arg Trp Arg Pro Leu Pro Gln Arg Glu Ser Gln
 20 25 30
 Gly Leu Met Gly Gly Asn Gly Arg Gly Thr Phe Thr Asp Arg Lys Ala
 35 40 45
 Gln Pro Gly Asp Phe Leu Gly Leu Leu Ala Arg Gly Thr Thr Pro Ser
 50 55 60

Pro Thr Thr Ala Ala Pro Ser Ser Arg Cys Trp Phe Arg Glu Gly Thr
 65 70 75 80
 Thr Met Tyr Ala Leu Tyr Ile Thr Val His Gly Tyr Phe Leu Ile Thr
 85 90 95
 Phe Leu Phe Gly Met Val Val Leu Ala Leu Val Val Trp Lys Ile Phe
 100 105 110
 Thr Leu Ser Arg Ala Thr Ala Val Lys Glu Arg Gly Lys Asn Arg Cys
 115 120 125
 Ser Pro Cys Trp Ala Ser Arg Ala Trp Trp Val
 130 135

<210> 43
 <211> 1650
 <212> DNA
 <213> Homo sapiens

<400> 43
 atggcgacgc ccaggggcct gggggccctg ctctgctcc tctgctccc gacctcaggt 60
 caggaaaagc ccaccgaagg gccaaagaac acctgacctg ggagcaacaa catgtacgac 120
 atcttcaact tgaatgacaa ggcttctgtc ttcaccaagt gcaggcagtc gggcagcgac 180
 tctgcaatg tggaaaactt gcagagatac tggctaaact acgaggcca tctgatgaag 240
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<210> 44
 <211> 549
 <212> PRT
 <213> Homo sapiens

<400> 44
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Leu	Cys	Phe	Thr	Lys	Cys	Arg	Gln	Ser	Gly	Ser	Asp	Ser	Cys	Asn	Val
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65					70					75					80
Glu	Gly	Leu	Thr	Gln	Lys	Val	Asn	Thr	Pro	Phe	Leu	Lys	Ala	Leu	Val
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Gln	Asn	Leu	Ser	Thr	Asn	Thr	Ala	Glu	Asp	Phe	Tyr	Phe	Ser	Leu	Glu
			100					105					110		
Pro	Ser	Gln	Val	Pro	Arg	Gln	Val	Met	Lys	Asp	Glu	Asp	Lys	Pro	Pro
		115					120					125			
Asp	Arg	Val	Arg	Leu	Pro	Lys	Ser	Leu	Phe	Arg	Ser	Leu	Pro	Gly	Asn
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Arg	Ser	Val	Val	Arg	Leu	Ala	Val	Thr	Ile	Leu	Asp	Ile	Gly	Pro	Gly
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Thr	Leu	Phe	Lys	Gly	Pro	Arg	Leu	Gly	Leu	Gly	Asp	Gly	Ser	Gly	Val
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Lys	Leu	Ala	Glu	Pro	Leu	Glu	Ile	Val	Phe	Ser	His	Gln	Arg	Pro	Pro
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Pro	Asn	Met	Thr	Leu	Thr	Cys	Val	Phe	Trp	Asp	Val	Thr	Lys	Gly	Thr
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Gly	Thr	Val	Cys	Cys	Cys	Asp	His	Leu	Thr	Phe	Phe	Ala	Leu	Leu	Leu
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Arg	Pro	Thr	Leu	Asp	Gln	Ser	Thr	Val	His	Ile	Leu	Thr	Arg	Ile	Ser
			260					265					270		
Gln	Ala	Gly	Cys	Gly	Val	Ser	Met	Ile	Phe	Leu	Ala	Phe	Thr	Ile	Ile
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	290					295					300				
Ala	Pro	Lys	Ile	His	Val	Ala	Leu	Gly	Gly	Ser	Leu	Phe	Leu	Leu	Asn
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Leu	Ala	Phe	Leu	Val	Asn	Val	Gly	Ser	Gly	Ser	Lys	Gly	Ser	Asp	Ala
			325						330					335	
Ala	Cys	Trp	Ala	Arg	Gly	Ala	Val	Phe	His	Tyr	Phe	Leu	Leu	Cys	Ala
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Phe	Thr	Trp	Met	Gly	Leu	Glu	Ala	Phe	His	Leu	Tyr	Leu	Leu	Ala	Val
		355					360					365			
Arg	Val	Phe	Asn	Thr	Tyr	Phe	Gly	His	Tyr	Phe	Leu	Lys	Leu	Ser	Leu
		370				375					380				
Val	Gly	Trp	Gly												

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Pro Leu Gly Leu Ser Thr Val Tyr Ile Phe Ala Leu Phe Asn Ser Leu
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Gln Gly Val Phe Ile Cys Cys Trp Phe Thr Ile Leu Tyr Leu Pro Ser
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Gln Ser Thr Thr Val Ser Ser Ser Thr Ala Arg Leu Asp Gln Ala His
530 535 540
Ser Ala Ser Gln Glu
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<210> 45
<211> 1020
<212> DNA
<213> Homo sapiens

<400> 45
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<210> 46
<211> 339
<212> PRT
<213> Homo sapiens

<400> 46
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Asp Trp Ser Ser Glu Gly Cys Ser Thr Glu Val Arg Pro Glu Gly Thr
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Val Cys Cys Cys Asp His Leu Thr Phe Phe Ala Leu Leu Leu Arg Pro
35 40 45
Thr Leu Asp Gln Ser Thr Val His Ile Leu Thr Arg Ile Ser Gln Ala
50 55 60
Gly Cys Gly Val Ser Met Ile Phe Leu Ala Phe Thr Ile Ile Leu Tyr
65 70 75 80
Ala Phe Leu Arg Leu Ser Arg Glu Arg Phe Lys Ser Glu Asp Ala Pro
85 90 95
Lys Ile His Val Ala Leu Gly Gly Ser Leu Phe Leu Leu Asn Leu Ala
100 105 110
Phe Leu Val Asn Val Gly Ser Gly Ser Lys Gly Ser Asp Ala Ala Cys

1203

<211> 400

<212> PRT

<213> Homo sapiens

<400> 48

31

Ser Ser Ser Thr Ala Arg Leu Asp Gln Ala His Ser Ala Ser Gln Glu
385 390 395 400

<210> 49
<211> 825
<212> DNA
<213> Homo sapiens

<400> 49
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<210> 50
<211> 274
<212> PRT
<213> Homo sapiens

<400> 50
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Ile His Val Ala Leu Gly Gly Ser Leu Phe Leu Leu Asn Leu Ala Phe
35 40 45
Leu Val Asn Val Gly Ser Gly Ser Lys Gly Ser Asp Ala Ala Cys Trp
50 55 60
Ala Arg Gly Ala Val Phe His Tyr Phe Leu Leu Cys Ala Phe Thr Trp
65 70 75 80
Met Gly Leu Glu Ala Phe His Leu Tyr Leu Leu Ala Val Arg Val Phe
85 90 95
Asn Thr Tyr Phe Gly His Tyr Phe Leu Lys Leu Ser Leu Val Gly Trp
100 105 110
Gly Leu Pro Ala Leu Met Val Ile Gly Thr Gly Ser Ala Asn Ser Tyr
115 120 125
Gly Leu Tyr Thr Ile Arg Asp Arg Glu Asn Arg Thr Ser Leu Glu Leu
130 135 140
Cys Trp Phe Arg Glu Gly Thr Thr Met Tyr Ala Leu Tyr Ile Thr Val
145 150 155 160
His Gly Tyr Phe Leu Ile Thr Phe Leu Phe Gly Met Val Val Leu Ala
165 170 175
Leu Val Val Trp Lys Ile Phe Thr Leu Ser Arg Ala Thr Ala Val Lys
180 185 190
Glu Arg Gly Lys Asn Arg Lys Lys Val Leu Thr Leu Leu Gly Leu Ser
195 200 205

165 170 175
 Ile Leu Tyr Leu Pro Ser Gln Ser Thr Thr Val Ser Ser Ser Thr Ala
 180 185 190
 Arg Leu Asp Gln Ala His Ser Ala Ser Gln Glu
 195 200

<210> 53
 <211> 4036
 <212> DNA
 <213> Homo sapiens

<400> 53
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